```
                                                                    Section 1
           (1)  1           10          20          30          40        52
ApoB  1-269 (1) --------------------MDPPRPALLALLA P LLLLLLLAGAR EEE LE
CD1d human  (1) MGCLLFLLLWALLQAWGSAEVPQRLFPLRCLQ S FANSSWTRTD LAW GE


                                                                    Section 2
           (53) 53        60          70          80          90       104
ApoB  1-269 (34) NV LVCPKD TRF HL K YN---------YEAE--S-SSGVPGTA SRS
CD1d human  (53) LQ HSWSND DTV SL P QGTFSDQQWETLQHIFRVYRSSFTRDVK FAK


                                                                    Section 3
          (105) 105   110         120         130         140        156
ApoB  1-269 (73) ATR NCK ELEVPQLCSFILK-- S---------------QCTLKEVYGFNP
CD1d human (105) MLR SYP ELQVSAGCEVHPGNA NNFFHVAFQGKDILSFQGTSWEPTQEAP


                                                                    Section 4
          (157) 157         170         180         190          208
ApoB  1-269 (108) EGKALLKKTKNS E AAAMSR ELKLAIPEGKQVFLYPEKDEPTYILNIKRG
CD1d human (157) LWVNLAIQVLNQ K TRETVQ LLNGTCPQFVSGLLESGKSE-----LKKQV


                                                                    Section 5
          (209) 209         220         230         240         250    260
ApoB  1-269 (160) IISALLVPPE EEAKQ LF DTVYGNCSTHFTVKTR GN----------A-
CD1d human (204) KPKAWLSRGP PGPGR LL CHVSGFYPKPVWVKWM GEQEQQGTQPGD LP


                                                                    Section 6
          (261) 261       270         280         290          300      312
ApoB  1-269 (201) -----T------ STERDLGQCDRFKPIRTGISP ALIKGM RPLSTLI S
CD1d human (256) NADETWYLRATL VAGEAAGLSCRVKHSSLEGQD VLYWGG YTSMGLI L


                                                                    Section 7
          (313) 313    320         330         343
ApoB  1-269 (242) QS---------CQ TLDAKRKHVAE CKEQ   SEQ ID NO: 11
CD1d human (308) VLACLLFLLIVG TSRFKRQTSYQ L---   SEQ ID NO: 12
```

Fig. 12

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                                                               ─────── Section 1
           (1) 1         10        20        30        40        52
ApoB 512-721  (1) ----------------------------------------------------K
CD1d human    (1) MGCLLFLLLWALLQAWGSAEVPQRLFPLRCLQISSFANSSWTRTDGLAWLGE

                                                               ─────── Section 2
          (53) 53      60        70        80        90        104
ApoB 512-721  (2) CVQS KPSLMIQKA  QALRKMEPKDKDQEVLLQTFLDDASPGD R AA LM
CD1d human   (53) LQTH WSNDSDTVR  KPWSQGTFSDQQWETLQHIFRVYRSSFT D KE AK

                                                               ─────── Section 3
         (105) 105   110       120       130       140        156
ApoB 512-721 (54)  RS--PSQAD NKIVQ LPWEQNE---------------QVKN VA HIAN
CD1d human  (105)  RLSYPLELQ SAGCE HPGNASNNFFHVAFQGKDILSFQGTS EP QEAP

                                                               ─────── Section 4
         (157) 157       170       180       190        208
ApoB 512-721 (89)  LNSEE D QDLKKLV EA KESQLPTVMDFRKFSRN------YQLYKSV L
CD1d human  (157)  WVNLA Q LNQDKWT ET QWLLNGTCPQFVSGLLESGKSELKKQVKPK W

                                                               ─────── Section 5
         (209) 209       220       230       240       250        260
ApoB 512-721(135) PSLDP S KIEGNL FDPNN LPK---ESM TTLTAFGFAS D EI LEG
CD1d human  (209) LSRGP P PGRLLL CHVSG YPKPVWVKW GEQEQQGTQP D PN DET

                                                               ─────── Section 6
         (261) 261       270       280       290       300        312
ApoB 512-721(184) KGFEPTL A FGKQ --------FFP SVNKALYW------------
CD1d human  (261) WYLRATL V AGEA GLSCRVKHSSL GQDIVLYWGGSYTSMGLIALAVLAC

                                                               ─────── Section 7
         (313) 313    320       335
ApoB 512-721(211) --------------------       SEQ ID NO: 13
CD1d human  (313) LLFLLIVGFTSRFKRQTSYQGVL    SEQ ID NO: 15
```

Fig. 13

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                                                                    Section 1
       (1) 1        10        20        30        40        52
ApoB 270-570 (1) --------FSYNNKYGMVAQVTQTLKLEDTPKINSRFFGEGT KMGLA EST
CD1d human   (1) MGCLLFLLLWALLQAWGSAEVPQRLFPLRCLQISSFANSSWT TDGLA LGE


                                                                    Section 2
      (53) 53       60        70        80        90        104
ApoB 270-570 (45) KSTSPPK AEAV  LQELKKLTIS Q IQRA LFN VTELRGLSDEAVTS
CD1d human   (53) LQTHSWS DSDT  LKPWSQGTFS Q WETL HIF YRSSFTRDVKEFAK


                                                                    Section 3
     (105) 105  110       120       130       140       156
ApoB 270-570 (97)  LPQLIE SSP LQALVQCG PQCSTHI Q LK VHANPLLIDVVTYLVAL
CD1d human  (105)  LRLSYP ELQ AGCEVHPG ASNNFFH A QG DILSFQGTSWEPTQEAP


                                                                    Section 4
     (157) 157       170       180       190       208
ApoB 270-570(149)  PEPSAQQ REIFNMAR QRSRATLYALSHAVNNYHK NP GTQEL DIAN
CD1d human  (157)  WVNLAIQ LNQDKWTR TVQWLLNGTCPQFVSGLLE GK ELKKQ KPKA


                                                                    Section 5
     (209) 209       220       230       240       250       260
ApoB 270-570(201) LMEQIQDD---------C GDEDYTYL LR IGNMGQTMEQLTPELKS ILK
CD1d human  (209) LSRGPSPGPGRLLLVCHV GFYPKPVW KW RGEQEQQGTQPGDILPN DET


                                                                    Section 6
     (261) 261       270       280       290       300       312
ApoB 270-570(244) CVQSTKPS IQKAAIQ LRKMEPKDKDQ  LQTFLDDASPGDKRLAAYLM
CD1d human  (261) WYLRATLD AGEAAGL CRVKHSSLEGQ  LYWGGSYTSMGLIALAVLAC


                                                                    Section 7
     (313) 313   320       335
ApoB 270-570(296) L RSPS------------------  SEQ ID NO: 14
CD1d human  (313) L FLLIVGFTSRFKRQTSYQGVL   SEQ ID NO: 16
```

Fig. 14